S. icy
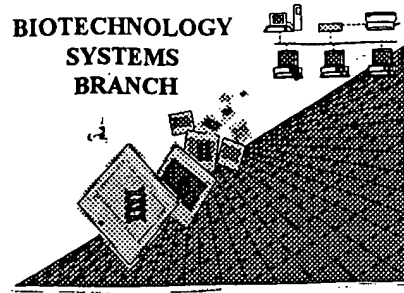folio

RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: _09/509,712 A_

Source: _1600 RUSH_

Date Processed by STIC: _11/7/2001_

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.
PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:
1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY
FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216.
PATENTIN 2.1 e-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax)
PATENTIN 3.0 e-mail help: patin3help@uspto.gov or phone 703-306-4119 (R. Wax)

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 3.0 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW:

Checker Version 3.0

The Checker Version 3.0 application is a state-of the-art Windows based software program employing a logical and intuitive user-interface to check whether a sequence listing is in compliance with format and content rules. Checker Version 3.0 works for sequence listings generated for the original version of 37 CFR §§1.821 – 1.825 effective October 1, 1990 (old rules) and the revised version (new rules) effective July 1, 1998 as well as World Intellectual Property Organization (WIPO) Standard ST.25.
Checker Version 3.0 replaces the previous DOS-based version of Checker, and is Y2K-compliant. Checker allows public users to check sequence listings in Computer Readable form (CRF) before submitting them to the United States Patent and Trademark Office (USPTO). Use of Checker prior to filing the sequence listing is expected to result in fewer errored sequence listings, thus saving time and money.

Checker Version 3.0 can be down loaded from the USPTO website at the following address:
http://www.uspto.gov/web/offices/pac/checker

1600

RAW SEQUENCE LISTING   DATE: 11/07/2001
PATENT APPLICATION: US/09/509,712A   TIME: 13:41:54

Input Set : N:\Crf3\09192001\I509712A.raw
Output Set: N:\CRF3\11072001\I509712A.raw

see pp. 1-5

Does Not Comply
Corrected Diskette Needed

```
 1 <110> APPLICANT: Rubin, Donald H.
 2       Organ, Edward L.
 3       DuBois, Raymond N.
 4 <120> TITLE OF INVENTION: Mammalian Genes Involved in Viral
 5       Infection and Tumor Suppression
 6 <130> FILE REFERENCE: 01123.0004
 7 <140> CURRENT APPLICATION NUMBER: US/09/509,712A
 8 <141> CURRENT FILING DATE: 2001-08-29
 9 <150> PRIOR APPLICATION NUMBER: PCT/US98/21276
10 <151> PRIOR FILING DATE: 1998-10-08
11 <150> PRIOR APPLICATION NUMBER: 60/062,021
12 <151> PRIOR FILING DATE: 1997-10-10
13 <160> NUMBER OF SEQ ID NOS: 127
14 <170> SOFTWARE: FastSEQ for Windows Version 4.0
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17 <211> LENGTH: 925
18 <212> TYPE: DNA
19 <213> ORGANISM: Rattus norvegicus
20 <221> NAME/KEY: misc_feature
21 <222> LOCATION: 1- 925
22 <223> OTHER INFORMATION: n = g, a, c or t(u)
23 <400> SEQUENCE: 1
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26       ggttgattgc tcttgaaagg ntttgaggtg naattcctcc gtnagtttga ccgtagtcgg     180
27       atntgaagag ggattgttna gcagncataa tttcattccc tgnacaccca gtaacnnttt     240
28       accgtcattt ggttgggaat tgatntcggg aggtancaan ggccacagtt atttattgtt     300
29       ncggaggatt gcaccaattn ggccggctgc ctctganatc tgtttctcat ccatgccggt     360
30       tcacccagac gaaagccgaa agcntcggga gtcctaactn tagtccntga aagtcattcc     420
31       cagctgcgta attgggctgt gcagagtccc agctcggtaa atatttgccc cgtgactgag     480
32       ctggagagaa tgctcctttc ttggtcctgg gcagctcttg gcagctcaca tgcactgttt     540
33       acctatcctc ccacattccc ccctgaggaa tcatcgtgcc tcggttccct taagtcctct     600
34       caacagaaaa caaggcagag tggaacgaag gaaagtgcgt ggccgttaga aagcctgtct     660
35       cgaatctgtc ccacgtgcct caggtagcgt tccaaacagc aaagattcta gtgaagaaaa     720
36       ataccgtccg gtcaattagt caggtggaca gagcaggacc cggtgtcttg gaagcctcgt     780
37       ccattcctct ggggaaggtg ggggggggcg tgtaatgcag ctctcaagaa gaaggtattt     840
38       ttgtttttcct ggagaaactg ccatcccagg agctgagagt ggatcagtag gaaggcctgt     900
39       gacaggaagc agggaggttc agcng                                         925
41 <210> SEQ ID NO: 2
42 <211> LENGTH: 554
43 <212> TYPE: DNA
44 <213> ORGANISM: Rattus norvegicus
45 <221> NAME/KEY: misc_feature
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47 <223> OTHER INFORMATION: n = g, a, c or t(u)
48 <400> SEQUENCE: 2
49       caagatngan ggggcggcgg ttcgnccaga gagcgggtag ggaagggaac gcgccggatg      60
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(global error)
insert this mandatory numeric identifier wherever <221>, <222>, <223> is shown

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/509,712A

DATE: 11/07/2001
TIME: 13:41:54

Input Set : N:\Crf3\09192001\I509712A.raw
Output Set: N:\CRF3\11072001\I509712A.raw

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W--> 51        ttgggaagag gaggaccgtg gttnataaat aacagaaagc ccagagggac gtanccatcc    180
W--> 52        gggatggaga gaggtaggga atccagntgt aagtcccaaa ctgccaccac cttcatnaga    240
W--> 53        actgcttcgt gtaaggtcac gcaccgggcc agctgtccng agtggcggtc ctggcgtgtt    300
W--> 54        aagttagcta aagtnactgc aactccgnct gtgcagactg ntcgtaaatt ctctctgtcc    360
W--> 55        gccaaattct ccctcctatt aaacttttca cttcctttca cttagtttcc tnacttcttt    420
W--> 56        caaacggaag ctgtaactga gcctgccacc cnganacntt gtggttgcca tttttatgct    480
W--> 57        aaagtaatcg tgtttttat gcctgtcaac tcccttttca tntaaagcag ggcntaccct    540
     58        attataactc tgcc                                                      554
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     61 <211> LENGTH: 891
     62 <212> TYPE: DNA
     63 <213> ORGANISM: Rattus norvegicus
     64 <221> NAME/KEY: misc_feature
     65 <222> LOCATION: 1- 825 878 (See below)
     66 <223> OTHER INFORMATION: n = g, a, c or t(u)
     67 <400> SEQUENCE: 3
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W--> 70        ntggtngggg ggggagggac ncccacagan tcatgnggtg gttngggngg ngggcatcgt    180
W--> 71        tnngatatta tcacattntg ngaanctatg tngggcttc ctttcngaca ggtggtggtt    240
W--> 72        nnacangngg atgtgtgctt ctttttcag cagtggtgga cccggattct aagaccctta    300
W--> 73        cngtaacaat gccctntttt cctaagccta accagtcctt tangaggant gctcttgggn    360
W--> 74        acccatgctg nntcacctag ccttggntca catnttnnac acaggaaaag gcagcatgtc    420
W--> 75        ttntnggagc tcagcttatt cccttcccnt cccatccagn atctccctgg gntggatgag    480
W--> 76        gtggatgacg catcttcaaa gcaccccacg tntcatggga tgtgcacagg agcttcgttg    540
W--> 77        gaaatgtgtt gcgcgaccag gcttgtgtag gaaacaacag actactcgaa attaaagtcn    600
W--> 78        taccttgcag ggttctcaga ggcttttacg cattaataaa catttgaatc ntaagaaggg    660
W--> 79        agcacagcat gtaatattnt tcaaattatc aggcnttgca accttcatta gtttctctta    720
W--> 80        cgcagctggg ngtggtggtg tgtacccttta atctcagcac tgaggaggca cngatatctc    780
W--> 81        catctctgtg acttccagac cggcntcgcc agagcaagtt ccaggccacc cagatgagat    840
W--> 82        gctcacagag gggacctttt tntgatgacc aacgnagnat gcaagtaagg a              891
     84 <210> SEQ ID NO: 4
     85 <211> LENGTH: 974
     86 <212> TYPE: DNA
     87 <213> ORGANISM: Rattus norvegicus
     88 <221> NAME/KEY: misc_feature
     89 <222> LOCATION: 1- 974
     90 <223> OTHER INFORMATION: n = g, a, c or t(u)
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W--> 94        nttattgccg ncntttcccc cccgctnttg cnccccttta cttngagant ngtgntncna    180
W--> 95        agatttnaag gttnttgccc ccccggcttt tnttcccctn nttttccccn nagntttaaa    240
W--> 96        accggtntgg gttncnantt nnttgnancc nccnattggg gtttccgntt accngggttt    300
W--> 97        ttccccatgn ccgttccctc caatnttgna cttcccnggt cngggtccna atnccnngna    360
W--> 98        acngntcnan ccttattgac aattaatttt tccttgngna ntctgncccc cngnantttg    420
W--> 99        gggttcttgg gngcagggcc ttttttttcnt tggnngcaan cncataaatn ttaccagntt    480
W--> 100       gattgctaag gaagtancca tggttgngaa ccccccccttn ttntctccca gatggaaccc    540
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W--> 101        aggattttgg aactgcagag gcttcagggt cttgggaagc ggaggcagnn aaagattgga    600
W--> 102        gtgcactgtc cttttgcaat atggggtttg cctgcctgct ggctcntctc ctgctntntc    660
W--> 103        agatggtgac tgaggctact tcngcaggac tnggaataat catgtccagg tggctgccct    720
W--> 104        tccgagcaga aagggacaga cgtggggcga tgaagttgct atcgtttntt tttttttctg    780
     105        cacagactgc aaagtgtgca gagggaggga ggctgtgcaa aaaaaaaaaa aaaaaaaaaa    840
     106        aaaaaaaaaa ccgaggacgc agaagttaga ctgctgaccc atttggtgca tgtgtgccca    900
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W--> 108        acgnnnctcc caga                                                      974
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    111 <211> LENGTH: 850
    112 <212> TYPE: DNA
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    115 <222> LOCATION: 1- 850
    116 <223> OTHER INFORMATION: n = g, a, c or t(u)
    117 <400> SEQUENCE: 5
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W--> 119        tgnggagntt ggngaccttn ttaccgnaan gagtgggaaa cgtttcctc cgggttnang    120
W--> 120        gttaggggga cccgnnggaa aatttttaaaa ccnngngggc tttttcgaat taaggggaaa    180
W--> 121        ngcggttttng gtnnntgaag ggcgggnggt tggagtcnna gtccagagtt gatttccacc    240
W--> 122        cacaaatntg ggaggtgncg gggaatgntg ncnttttctt gngatgaggg ntgccgtncc    300
W--> 123        ggantaacag ngnttgcntt gtntngcaaa acgaagagtn tcctgnttgg aataggngtt    360
W--> 124        cngttcgang ganccagatt tangngntgg agnaaggatt nggcagataa angcntgaga    420
W--> 125        natgnancnt ggancaggtc nggncnnagn ntacagatga tgnncccana canganataa    480
W--> 126        ntncagatca cagtcgtacc cgnggctggg ccatgaanag ggcatcccca gacnnacaca    540
W--> 127        ngccttnana antgntcaga gaaccancag tggntanggg ntgcccnnnn naccagggaa    600
W--> 128        gacccggggc gtgncggata ttgacacanc agatnncatt tggggncggt tcgagggttn    660
W--> 129        atgntcnccg agtacnagan angatcntcc aacccggaat ncggtgctcc ngtcgtccga    720
W--> 130        tgnaatgagt cgnccggnaa cctcatatcc aagaaacnat acagcagtgg nntccgagtc    780
W--> 131        tcgtatantc nttgcggggng gaggctatnt tcagaggnca agattaccgt tagcgggana    840
W--> 132        aagtngaana                                                           850
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    135 <211> LENGTH: 531
    136 <212> TYPE: DNA
    137 <213> ORGANISM: Rattus norvegicus
    138 <221> NAME/KEY: misc_feature
    139 <222> LOCATION: 1- 531
    140 <223> OTHER INFORMATION: n = g, a, c or t(u)
    141 <400> SEQUENCE: 6
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W--> 143        ngtctnntgt ctgtgtngtg cccctgtccn catctctcac nccagggaga gagatgtgag    120
W--> 144        ananacatca gagatctctn gnacagtgtt tcacaagagt ctatcncana gagcacatct    180
W--> 145        gcccgggggng anacacaact ctaaatgtgt ctcanntgat ctctctnttg tgtctctnac    240
W--> 146        atatgnggac atgctctcag agtatnggnt ctcttgngcn cttntgcaca cacacacaca    300
W--> 147        cacacacaca cacacacaca cacncttctc tctggcacag ggntatggca nagcacatnt    360
W--> 148        tnngagntca nagctntata tgagtgtgtg gcgaaaggng tnatnanann gacnnccca    420
W--> 149        gcnnatatag gggggngnnc tctnggcctc tcttnggnaa tntgngggng agtctgcnca    480
W--> 150        cacaggcgct cnnacccanc nnnttggggc cccccaggng ttttcncccc c             531
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Input Set : N:\Crf3\09192001\I509712A.raw
Output Set: N:\CRF3\11072001\I509712A.raw

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        154 <212> TYPE: DNA
        155 <213> ORGANISM: Rattus norvegicus           >(2207
        156 <221> NAME/KEY: misc_feature
        157 <222> LOCATION: 1- 572
        158 <223> OTHER INFORMATION: n = g, a, c or t(u)
        159 <400> SEQUENCE: 7
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W-->    161     gagacancgg nnacacagag gttttgngnn tattgngagt ctctgcgcac nccananttt     120
W-->    162     aaccncgggg nctcntgttt tattttaaaa aaaaagagtc ncatgtntat ttctctnatg     180
W-->    163     tgaaaatcnc attcanagtt ntggggtttc ccntgaggag anatagagtt tcacactctt     240
W-->    164     ctctccgagg ggtcntcnca tgtntctccc caatgtgngn ggnacacaca tgnggccccn     300
W-->    165     aggggggtgng ctctctctgc ncagggcncc ccccaanang tagaganaca ntgtggtgtt     360
W-->    166     tcacaacaca attcncgaga nattntgttc cncantggnn gtctnagntc ncatgttgtg     420
W-->    167     gngacangtt agnncncccc atnttcnccc ccctttcaca ctgccccnag agagagaaan     480
W-->    168     tctnggcccc ctctanannt nttttaaat cnccccnnac cacaggtntt cccagggtat     540
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        173 <212> TYPE: DNA
        174 <213> ORGANISM: Rattus norvegicus           >(2207
        175 <221> NAME/KEY: misc_feature
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        177 <223> OTHER INFORMATION: n = g, a, c or t(u)
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W-->    180     gcgagaanac tctgtnnant ngtctccccc cncnccnaca gngtganant caaaacctct     120
W-->    181     agagcccccc agaaanccccc tntctcaaan aaagagaaag agaagancga gnagnagaga     180
W-->    182     gananagaga gagagagtgt gganctntnt cctcngancc ccannnanan ngtgnggcnc     240
W-->    183     actcncnngt gnngngnacc ccngggggatt tncgcgtgtc cccttgngct ctgtntanga     300
W-->    184     gananatatg tntagtctct ctntcgcccc ctccgntgtc acgtgtgcgg ggcccnngag     360
W-->    185     acacagacac ntctctcang gggaacacat anngactcnc acntgtgttt atattcnccc     420
W-->    186     ctcccnctca cacanacaca cacacagnag atattnngct actctctctc tgtcacaggg     480
W-->    187     gtacanattt antctnggcc anacccctct cngaagngng ggcanngtaa accccgcccc     540
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W-->    189     cccttngnac cccccctntna accctctntt tgaacncaac ncaccntccc cnttttctcg     660
W-->    190     gggnnggncc ngcnccncct ctcncaaaaa aaattnnaan ttngtcccct ncccnttnt     720
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W-->    192     ttcccctttt ccnggggacc cccgnnttcc tttttaaaaa aaaanaaccc tttctccctt     840
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        202 <223> OTHER INFORMATION: n = g, a, c or t(u)
        203 <400> SEQUENCE: 9
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W--> 211          ccaccncnan gacacaaana naaggngtnn anaaccccan aaaaactnng ntntcngntt         480
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W--> 214          tntncttacc cctctcccct ctctgtgtgt ngannatatc ngntctagat gtcctnaccc         660
W--> 215          tccccaaacc tttctcnggn agagacntct ctntnttttt ccccncttc catttgaaan          720
W--> 216          anangagaag gnccaaaaag gngggngtct tctcgggaat ncnccctttt ggcccccccaa         780
W--> 217          cctgggtttt tttcccccctt cctttttaatn antttttcna nacaaanctt tnngngtttn         840
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     223 <212> TYPE: DNA
     224 <213> ORGANISM: Rattus norvegicus
     225 <221> NAME/KEY: misc_feature
     226 <222> LOCATION: 1- 400
     227 <223> OTHER INFORMATION: n = g, a, c or t(u)
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     230          acagtttatg aatattatta catatttcac aagactttat attgttataa tatgcctcat         120
     231          gtgagatata tgtgattctg tggtggtgtt ctcagagggg gtttgggtta ttggggataa         180
     232          tagtttgccc ctcgcggggt ctatatttat atatgtgaca caatatatta gagagatttt         240
     233          tggttatata tatttccctt cgcgggggtg gagatttatc acaggggag agctttttccc          300
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     238 <211> LENGTH: 880
     239 <212> TYPE: DNA
     240 <213> ORGANISM: Rattus norvegicus
     241 <221> NAME/KEY: misc_feature
     242 <222> LOCATION: 1- 880
     243 <223> OTHER INFORMATION: n = g, a, c or t(u)
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W--> 248          aatgaaggtn ttccggaggg ttgccttcca atcccagatg gatttgagtt tcaggatcaa         240
W--> 249          ttcagttacc gntgaccatc caccnncctc cngtataatc attngatgag gatgaatggt         300
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     254          ggagaaaatt ccagtccagt gagtcatggg cagggggagg ggtggggtaa gggcagtggc         600
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VERIFICATION SUMMARY　　　　　　　　DATE: 11/07/2001
PATENT APPLICATION:　US/09/509,712A　　TIME: 13:41:55

Input Set : N:\Crf3\09192001\I509712A.raw
Output Set: N:\CRF3\11072001\I509712A.raw

```
L:7 M:270 C: Current Application Number differs, Wrong Format
L:24 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:1
L:24 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1
L:25 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:1
L:25 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1
L:26 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:1
L:26 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1
L:27 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:1
L:27 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1
L:28 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:1
L:28 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1
L:29 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:1
L:29 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1
L:30 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:1
L:30 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1
L:39 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:1
L:39 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1
L:49 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:2
L:49 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2
L:50 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:2
L:50 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2
L:51 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:2
L:51 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2
L:52 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:2
L:52 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2
L:53 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:2
L:53 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2
L:54 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:2
L:54 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2
L:55 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:2
L:55 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2
L:56 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:2
L:56 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2
L:57 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:2
L:57 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2
L:68 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:3
L:68 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3
L:69 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:3
L:69 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3
L:70 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:3
L:70 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3
L:71 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:3
L:71 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3
L:72 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:3
L:72 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3
L:73 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:3
L:73 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3
L:74 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:3
```

VERIFICATION SUMMARY   DATE: 11/07/2001
PATENT APPLICATION: US/09/509,712A   TIME: 13:41:55

Input Set : N:\Crf3\09192001\I509712A.raw
Output Set: N:\CRF3\11072001\I509712A.raw

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L:74  M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3
L:75  M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:3
L:75  M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3
L:76  M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:3
L:76  M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3
L:77  M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:3
L:77  M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3
L:78  M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:3
L:78  M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3
L:79  M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:3
L:79  M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3
L:80  M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:3
L:80  M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3
L:81  M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:3
L:81  M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3
L:82  M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:3
L:82  M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3
L:92  M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:4
L:92  M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4
L:93  M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:4
L:93  M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4
L:94  M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:4
L:94  M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4
L:95  M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:4
L:95  M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4
L:96  M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:4
L:96  M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4
L:97  M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:4
L:97  M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4
L:98  M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:4
L:98  M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4
L:99  M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:4
L:99  M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4
L:100 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:4
L:100 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4
L:101 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:4
L:101 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4
L:102 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:4
L:102 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4
L:103 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:4
L:103 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4
L:104 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:4
L:104 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4
L:107 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:4
L:107 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4
L:108 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:4
L:108 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4
L:118 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:5
L:118 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5
```

VERIFICATION SUMMARY　　　　　　　　　DATE: 11/07/2001
PATENT APPLICATION:　US/09/509,712A　　　TIME: 13:41:55

Input Set : **N:\Crf3\09192001\I509712A.raw**
Output Set: **N:\CRF3\11072001\I509712A.raw**

L:119 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:5
L:119 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5
L:120 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:5
L:120 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5